

# Potassium morpholine-4-carbodithioate monohydrate

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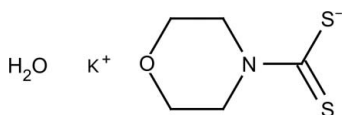
Received 30 May 2012; accepted 28 June 2012

Key indicators: single-crystal X-ray study;  $T = 290$  K; mean  $\sigma(\text{C}—\text{C}) = 0.005$  Å;  
 $R$  factor = 0.047;  $wR$  factor = 0.130; data-to-parameter ratio = 24.7.

In the ionic title compound,  $\text{K}^+ \cdot \text{C}_5\text{H}_8\text{NOS}_2^- \cdot \text{H}_2\text{O}$ , the morpholine ring of the morpholine-4-carbodithioate anion has a chair conformation. The potassium cation is coordinated by four S and four O atoms in a bipyramidal reversed geometry. In the crystal, the three components are linked, generating infinite two-dimensional networks that lie parallel to the  $bc$  plane. These layers are linked *via*  $\text{O}—\text{H} \cdots \text{S}$  hydrogen bonds, forming a three-dimensional structure.

## Related literature

For the crystal structures of similar compounds, see: Oskarsson *et al.* (1979); Albertsson *et al.* (1980); Ymén (1982); Mafud & Gambardella (2011*a,b*); Mafud *et al.* (2011). For puckering parameters, see: Cremer & Pople (1975).



## Experimental

### Crystal data

$\text{K}^+ \cdot \text{C}_5\text{H}_8\text{NOS}_2^- \cdot \text{H}_2\text{O}$   
 $M_r = 219.36$   
Monoclinic,  $P2_1/c$   
 $a = 6.7235$  (10) Å  
 $b = 17.260$  (4) Å  
 $c = 8.1904$  (10) Å  
 $\beta = 108.994$  (10)°  
 $V = 898.7$  (3) Å<sup>3</sup>  
 $Z = 4$   
Mo  $K\alpha$  radiation  
 $\mu = 1.01$  mm<sup>-1</sup>  
 $T = 290$  K  
 $0.45 \times 0.30 \times 0.20$  mm

### Data collection

Enraf–Nonius TurboCAD-4  
diffractometer  
Absorption correction: refined from  
 $\Delta F$  (Walker & Stuart, 1983)  
 $T_{\min} = 0.512$ ,  $T_{\max} = 0.818$   
2779 measured reflections  
2618 independent reflections  
1615 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.027$   
3 standard reflections every 120 min  
intensity decay: 10%

## Refinement

$R[F^2 > 2\sigma(F^2)] = 0.047$   
 $wR(F^2) = 0.130$   
 $S = 1.01$   
2618 reflections  
106 parameters  
3 restraints

H atoms treated by a mixture of  
independent and constrained  
refinement  
 $\Delta\rho_{\max} = 0.50$  e Å<sup>-3</sup>  
 $\Delta\rho_{\min} = -0.61$  e Å<sup>-3</sup>

Table 1

Hydrogen-bond geometry (Å, °).

$D—H \cdots A$	$D—H$	$H \cdots A$	$D \cdots A$	$D—H \cdots A$
$\text{O2}—\text{H1O} \cdots \text{S1}^{\text{i}}$	0.86 (4)	2.45 (4)	3.219 (3)	149 (3)
$\text{O2}—\text{H2O} \cdots \text{S1}^{\text{ii}}$	0.85 (3)	2.87 (5)	3.462 (3)	129 (4)

Symmetry codes: (i)  $-x - 1, y + \frac{1}{2}, -z + \frac{1}{2}$ ; (ii)  $-x - 1, -y, -z + 1$ .

Data collection: *CAD-4 EXPRESS* (Enraf–Nonius, 1994); cell refinement: *CAD-4 EXPRESS*; data reduction: *XCAD4* (Harms & Wocadlo, 1995); program(s) used to solve structure: *SIR92* (Altomare *et al.*, 1994); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

The author is grateful to the CNPq, National Counsel of Technological and Scientific Development, for supporting this study.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: SU2446).

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## supplementary materials

*Acta Cryst.* (2012). E68, m1025 [doi:10.1107/S1600536812029613]

**Potassium morpholine-4-carbodithioate monohydrate**

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**Comment**

The title compound, Fig. 1, is composed of a morpholinedithiocarbamate anion in contact with a potassium cation, which in turn is linked with a water molecule of crystallization. The crystal structure of similar compounds, for example Sodium 1-*R*-carbodithioate dihydrate, have been reported (Oskarsson *et al.*, 1979; Albertsson *et al.*, 1980; Ymén, 1982; Mafud & Gambardella *et al.*, 2011*a,b*).

The six-membered morpholine ring has a chair conformation with Puckering parameters [Cremer & Pople, 1975)]  $Q = 0.548(3) \text{ \AA}$ ,  $\theta = 173.3(3)^\circ$ ,  $\phi_2 = 2.6(3,4)^\circ$ .

In the crystal, a polymeric structure is built by coordination of the potassium cation to four sulfur [ $K \cdots S = 3.2670(13) - 3.3797(14) \text{ \AA}$ ] and four oxygen [ $K \cdots O = 2.828(3) - 3.007(3) \text{ \AA}$ ] atoms, with a bi-pyramidal reversed geometry. This configuration generates close packed layers which remain cohesive in crystal stacking by van der Waals interactions. The distances of these contacts are slightly less than the sum of the van der Waals radii.

The crystal packing gives rise to a supramolecular structure, whose infinite two-dimensional network lies parallel to the *bc* plane (Fig. 2). These layers are linked via O-H $\cdots$ S hydrogen bonds (Table 1) to form a three-dimensional structure.

**Experimental**

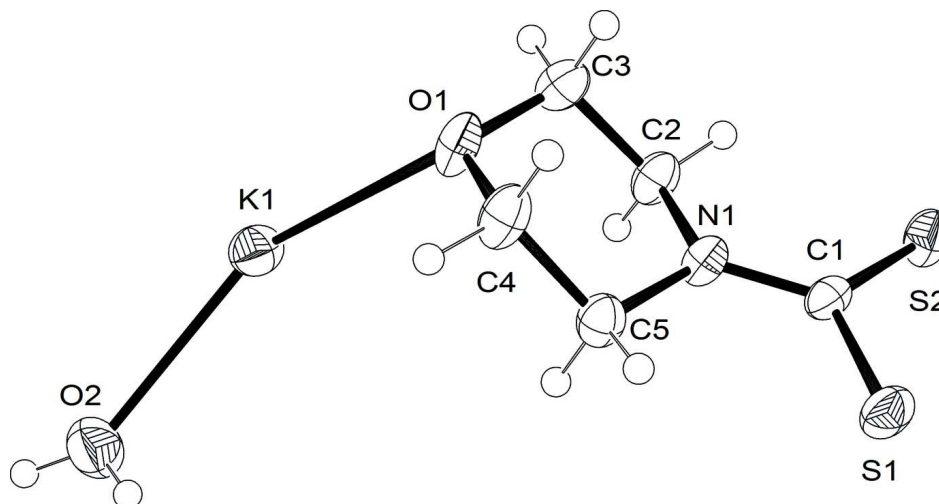
The potassium salts of DTC were prepared by direct reaction between amine and carbon disulfide (CS<sub>2</sub>) in the presence of a stoichiometric amount of potassium hydroxide in ethanol/water 1:1 (*v:v*). The reaction mixture was placed in the freezer for 12 h and then filtered through a Büchner funnel, washed with cold ether and the product recrystallized in an ethanol-water mixture 1:1 (*v:v*). The obtained solid was recrystallized from ethanol-water 1:1 (*v:v*) and dried in a vacuum oven at 323 K for 8 h. Colourless crystals, suitable for X-ray diffraction analysis, were obtained. On heating they sublimed and decomposed.

**Refinement**

The H-atom positions of the water molecule were located in a difference Fourier map and were refined with  $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{O})$ ; O—H = 0.86(4) and 0.85(3) Å. The C-bound H-atoms of the anion were included in calculated positions and treated as riding atoms: C—H = 0.97 Å, with  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{parent C-atom})$ .

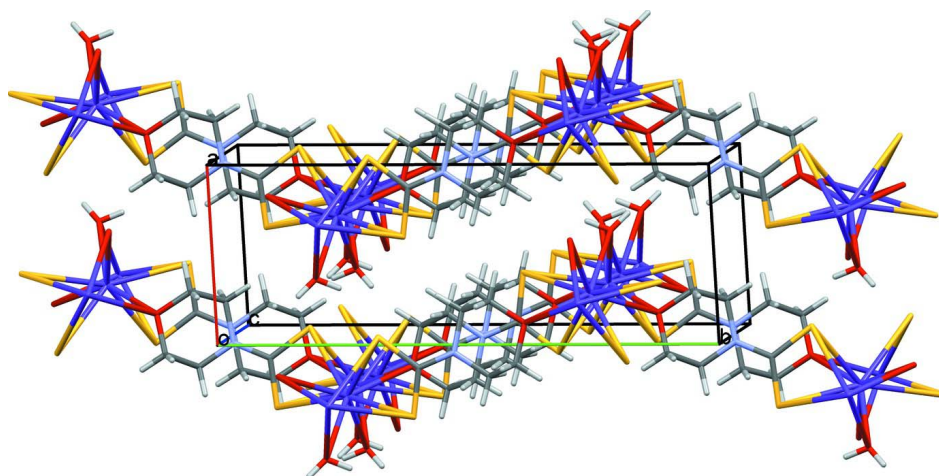
**Computing details**

Data collection: *CAD-4 EXPRESS* (Enraf–Nonius, 1994); cell refinement: *CAD-4 EXPRESS* (Enraf–Nonius, 1994); data reduction: *XCAD4* (Harms & Wocadlo, 1995); program(s) used to solve structure: *SIR92* (Altomare *et al.*, 1994); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997); software used to prepare material for publication: *WinGX* (Farrugia, 1999).



**Figure 1**

A view of the molecular structure of the asymmetric unit of the title compound, with the numbering scheme. The displacement ellipsoids are drawn at the 50% probability level.



**Figure 2**

The view along the *c* axis of the crystal packing of the title compound.

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#### Crystal data

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$M_r = 219.36$

Monoclinic,  $P2_1/c$

Hall symbol:  $-P\ 2_1/c$

$a = 6.7235\ (10)\ \text{\AA}$

$b = 17.260\ (4)\ \text{\AA}$

$c = 8.1904\ (10)\ \text{\AA}$

$\beta = 108.994\ (10)^\circ$

$V = 898.7\ (3)\ \text{\AA}^3$

$Z = 4$

$F(000) = 456$

$D_x = 1.621\ \text{Mg m}^{-3}$

Mo  $K\alpha$  radiation,  $\lambda = 0.71073\ \text{\AA}$

Cell parameters from 16 reflections

$\theta = 9.8\text{--}18.3^\circ$

$\mu = 1.01\ \text{mm}^{-1}$

$T = 290\ \text{K}$

Prism, colourless

$0.45 \times 0.3 \times 0.2\ \text{mm}$

### Data collection

Enraf–Nonius TurboCAD-4  
diffractometer  
Radiation source: Enraf Nonius FR590  
Graphite monochromator  
non-profiled  $\omega/2\theta$  scans  
Absorption correction: part of the refinement  
model ( $\Delta F$ )  
(Walker & Stuart, 1983)  
 $T_{\min} = 0.512$ ,  $T_{\max} = 0.818$   
2779 measured reflections

2618 independent reflections  
1615 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.027$   
 $\theta_{\max} = 30.0^\circ$ ,  $\theta_{\min} = 2.4^\circ$   
 $h = -9 \rightarrow 8$   
 $k = 0 \rightarrow 24$   
 $l = 0 \rightarrow 11$   
3 standard reflections every 120 min  
intensity decay: 10%

### Refinement

Refinement on  $F^2$   
Least-squares matrix: full  
 $R[F^2 > 2\sigma(F^2)] = 0.047$   
 $wR(F^2) = 0.130$   
 $S = 1.01$   
2618 reflections  
106 parameters  
3 restraints  
Primary atom site location: structure-invariant  
direct methods

Secondary atom site location: difference Fourier  
map  
Hydrogen site location: inferred from  
neighbouring sites  
H atoms treated by a mixture of independent  
and constrained refinement  
 $w = 1/[\sigma^2(F_o^2) + (0.0644P)^2]$   
where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\max} < 0.001$   
 $\Delta\rho_{\max} = 0.50 \text{ e } \text{\AA}^{-3}$   
 $\Delta\rho_{\min} = -0.61 \text{ e } \text{\AA}^{-3}$

### Special details

**Geometry.** Bond distances, angles etc. have been calculated using the rounded fractional coordinates. All su's are estimated from the variances of the (full) variance-covariance matrix. The cell esds are taken into account in the estimation of distances, angles and torsion angles

**Refinement.** Refinement of  $F^2$  against ALL reflections. The weighted  $R$ -factor  $wR$  and goodness of fit  $S$  are based on  $F^2$ , conventional  $R$ -factors  $R$  are based on  $F$ , with  $F$  set to zero for negative  $F^2$ . The threshold expression of  $F^2 > \sigma(F^2)$  is used only for calculating  $R$ -factors(gt) etc. and is not relevant to the choice of reflections for refinement.  $R$ -factors based on  $F^2$  are statistically about twice as large as those based on  $F$ , and  $R$ -factors based on ALL data will be even larger.

### Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters ( $\text{\AA}^2$ )

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
K1	−0.28085 (12)	0.24637 (4)	0.48217 (9)	0.0366 (2)
S1	−0.06278 (13)	−0.16683 (4)	0.69255 (11)	0.0320 (2)
S2	0.36659 (13)	−0.09853 (4)	0.80129 (12)	0.0336 (3)
O1	−0.1501 (4)	0.13219 (12)	0.7743 (3)	0.0366 (7)
O2	−0.5955 (4)	0.27515 (16)	0.1617 (4)	0.0496 (9)
N1	0.0172 (4)	−0.01514 (13)	0.7295 (3)	0.0261 (7)
C1	0.1003 (5)	−0.08662 (15)	0.7399 (4)	0.0240 (8)
C2	0.1460 (5)	0.05521 (16)	0.7583 (4)	0.0293 (9)
C3	0.0660 (5)	0.11438 (18)	0.8574 (4)	0.0334 (10)
C4	−0.2719 (5)	0.06303 (18)	0.7624 (5)	0.0348 (10)
C5	−0.2096 (4)	0.00131 (17)	0.6581 (4)	0.0289 (8)
H1O	−0.682 (6)	0.3070 (18)	0.093 (5)	0.0740*
H2A	0.14220	0.07690	0.64810	0.0350*
H2B	0.29080	0.04230	0.82250	0.0350*
H2O	−0.645 (7)	0.2294 (12)	0.154 (6)	0.0740*

H3A	0.08450	0.09460	0.97230	0.0400*
H3B	0.14850	0.16140	0.86940	0.0400*
H4A	−0.41980	0.07520	0.70910	0.0420*
H4B	−0.25230	0.04350	0.87770	0.0420*
H5A	−0.28850	−0.04570	0.65890	0.0350*
H5B	−0.24380	0.01850	0.53940	0.0350*

*Atomic displacement parameters (Å<sup>2</sup>)*

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
K1	0.0408 (4)	0.0356 (4)	0.0341 (4)	−0.0006 (3)	0.0130 (3)	−0.0026 (3)
S1	0.0383 (4)	0.0207 (3)	0.0366 (4)	−0.0060 (3)	0.0116 (3)	−0.0032 (3)
S2	0.0297 (4)	0.0270 (4)	0.0454 (5)	0.0033 (3)	0.0141 (3)	−0.0004 (3)
O1	0.0370 (12)	0.0220 (10)	0.0551 (15)	0.0040 (9)	0.0210 (11)	−0.0031 (10)
O2	0.0444 (15)	0.0393 (14)	0.0546 (17)	0.0006 (12)	0.0018 (13)	0.0048 (13)
N1	0.0270 (12)	0.0204 (11)	0.0339 (14)	−0.0009 (9)	0.0142 (10)	−0.0035 (10)
C1	0.0313 (14)	0.0203 (13)	0.0231 (14)	−0.0011 (11)	0.0124 (11)	−0.0031 (11)
C2	0.0310 (15)	0.0205 (13)	0.0400 (18)	−0.0044 (11)	0.0164 (14)	−0.0021 (12)
C3	0.0356 (17)	0.0278 (15)	0.0381 (18)	−0.0040 (12)	0.0140 (14)	−0.0072 (13)
C4	0.0326 (16)	0.0267 (15)	0.051 (2)	0.0004 (12)	0.0218 (15)	−0.0025 (14)
C5	0.0267 (14)	0.0255 (14)	0.0349 (16)	−0.0005 (11)	0.0108 (13)	−0.0004 (12)

*Geometric parameters (Å, °)*

K1—O1	3.002 (2)	N1—C2	1.465 (4)
K1—O2	2.828 (3)	N1—C1	1.346 (4)
K1—S1 <sup>i</sup>	3.2670 (13)	N1—C5	1.472 (4)
K1—S2 <sup>i</sup>	3.3630 (13)	C2—C3	1.508 (4)
K1—S1 <sup>ii</sup>	3.3797 (14)	C4—C5	1.508 (5)
K1—S2 <sup>ii</sup>	3.3708 (13)	C2—H2A	0.9700
K1—O1 <sup>iii</sup>	3.007 (3)	C2—H2B	0.9700
K1—O2 <sup>iv</sup>	2.967 (3)	C3—H3A	0.9700
S1—C1	1.730 (3)	C3—H3B	0.9700
S2—C1	1.707 (4)	C4—H4A	0.9700
O1—C3	1.423 (4)	C4—H4B	0.9700
O1—C4	1.433 (4)	C5—H5A	0.9700
O2—H2O	0.85 (3)	C5—H5B	0.9700
O2—H1O	0.86 (4)		
O1—K1—O2	142.34 (8)	K1 <sup>iv</sup> —O1—C3	108.72 (17)
S1 <sup>i</sup> —K1—O1	72.85 (5)	K1 <sup>iv</sup> —O1—C4	110.7 (2)
S2 <sup>i</sup> —K1—O1	99.11 (5)	K1—O2—K1 <sup>iii</sup>	89.96 (8)
S1 <sup>ii</sup> —K1—O1	90.44 (6)	K1 <sup>iii</sup> —O2—H2O	100 (3)
S2 <sup>ii</sup> —K1—O1	89.55 (5)	K1—O2—H2O	94 (3)
O1—K1—O1 <sup>iii</sup>	147.70 (8)	H1O—O2—H2O	112 (4)
O1—K1—O2 <sup>iv</sup>	66.04 (8)	K1—O2—H1O	149 (2)
S1 <sup>i</sup> —K1—O2	142.49 (6)	K1 <sup>iii</sup> —O2—H1O	101 (3)
S2 <sup>i</sup> —K1—O2	98.33 (6)	C1—N1—C5	123.9 (3)
S1 <sup>ii</sup> —K1—O2	94.95 (6)	C2—N1—C5	112.7 (2)
S2 <sup>ii</sup> —K1—O2	65.49 (6)	C1—N1—C2	122.6 (3)

O1 <sup>iii</sup> —K1—O2	67.68 (8)	S1—C1—N1	120.0 (3)
O2—K1—O2 <sup>iv</sup>	92.48 (9)	S2—C1—N1	120.2 (2)
S1 <sup>i</sup> —K1—S2 <sup>i</sup>	53.28 (3)	S1—C1—S2	119.81 (16)
S1 <sup>i</sup> —K1—S1 <sup>ii</sup>	97.59 (3)	N1—C2—C3	110.6 (3)
S1 <sup>i</sup> —K1—S2 <sup>ii</sup>	145.67 (4)	O1—C3—C2	112.1 (3)
S1 <sup>i</sup> —K1—O1 <sup>iii</sup>	83.26 (5)	O1—C4—C5	111.7 (3)
S1 <sup>i</sup> —K1—O2 <sup>iv</sup>	94.72 (6)	N1—C5—C4	110.7 (3)
S1 <sup>ii</sup> —K1—S2 <sup>i</sup>	143.59 (3)	N1—C2—H2A	110.00
S2 <sup>i</sup> —K1—S2 <sup>ii</sup>	161.01 (4)	N1—C2—H2B	109.00
S2 <sup>i</sup> —K1—O1 <sup>iii</sup>	82.85 (5)	C3—C2—H2A	109.00
S2 <sup>i</sup> —K1—O2 <sup>iv</sup>	64.28 (6)	C3—C2—H2B	110.00
S1 <sup>ii</sup> —K1—S2 <sup>ii</sup>	52.27 (3)	H2A—C2—H2B	108.00
S1 <sup>ii</sup> —K1—O1 <sup>iii</sup>	71.17 (5)	O1—C3—H3A	109.00
S1 <sup>ii</sup> —K1—O2 <sup>iv</sup>	148.77 (6)	O1—C3—H3B	109.00
S2 <sup>ii</sup> —K1—O1 <sup>iii</sup>	98.84 (5)	C2—C3—H3A	109.00
S2 <sup>ii</sup> —K1—O2 <sup>iv</sup>	104.89 (6)	C2—C3—H3B	109.00
O1 <sup>iii</sup> —K1—O2 <sup>iv</sup>	139.03 (8)	H3A—C3—H3B	108.00
K1 <sup>v</sup> —S1—C1	87.47 (11)	O1—C4—H4A	109.00
K1 <sup>ii</sup> —S1—C1	87.09 (11)	O1—C4—H4B	109.00
K1 <sup>v</sup> —S1—K1 <sup>ii</sup>	76.09 (3)	C5—C4—H4A	109.00
K1 <sup>v</sup> —S2—C1	84.71 (10)	C5—C4—H4B	109.00
K1 <sup>ii</sup> —S2—C1	87.73 (10)	H4A—C4—H4B	108.00
K1 <sup>v</sup> —S2—K1 <sup>ii</sup>	74.95 (2)	N1—C5—H5A	110.00
K1—O1—C3	120.77 (19)	N1—C5—H5B	110.00
K1—O1—C4	118.7 (2)	C4—C5—H5A	109.00
K1—O1—K1 <sup>iv</sup>	85.98 (6)	C4—C5—H5B	109.00
C3—O1—C4	108.9 (2)	H5A—C5—H5B	108.00
O2—K1—O1—C3	142.4 (2)	O2—K1—S1 <sup>ii</sup> —C1 <sup>ii</sup>	−72.98 (12)
O2—K1—O1—C4	3.2 (3)	O2—K1—S1 <sup>ii</sup> —K1 <sup>iii</sup>	15.13 (6)
O2—K1—O1—K1 <sup>iv</sup>	−108.20 (12)	O1—K1—S2 <sup>ii</sup> —C1 <sup>ii</sup>	−71.26 (12)
S1 <sup>i</sup> —K1—O1—C3	−54.13 (19)	O1—K1—S2 <sup>ii</sup> —K1 <sup>iii</sup>	−156.39 (6)
S1 <sup>i</sup> —K1—O1—C4	166.7 (2)	O2—K1—S2 <sup>ii</sup> —C1 <sup>ii</sup>	137.86 (12)
S1 <sup>i</sup> —K1—O1—K1 <sup>iv</sup>	55.30 (5)	O2—K1—S2 <sup>ii</sup> —K1 <sup>iii</sup>	52.73 (7)
S2 <sup>i</sup> —K1—O1—C3	−101.0 (2)	O1—K1—O1 <sup>iii</sup> —K1 <sup>iii</sup>	111.73 (12)
S2 <sup>i</sup> —K1—O1—C4	119.8 (2)	O1—K1—O1 <sup>iii</sup> —C3 <sup>iii</sup>	−9.5 (2)
S2 <sup>i</sup> —K1—O1—K1 <sup>iv</sup>	8.40 (6)	O1—K1—O1 <sup>iii</sup> —C4 <sup>iii</sup>	−129.1 (2)
S1 <sup>ii</sup> —K1—O1—C3	43.7 (2)	O2—K1—O1 <sup>iii</sup> —K1 <sup>iii</sup>	−50.45 (8)
S1 <sup>ii</sup> —K1—O1—C4	−95.5 (2)	O2—K1—O1 <sup>iii</sup> —C3 <sup>iii</sup>	−171.6 (2)
S1 <sup>ii</sup> —K1—O1—K1 <sup>iv</sup>	153.11 (5)	O2—K1—O1 <sup>iii</sup> —C4 <sup>iii</sup>	68.8 (2)
S2 <sup>ii</sup> —K1—O1—C3	96.0 (2)	O1—K1—O2 <sup>iv</sup> —K1 <sup>iv</sup>	52.11 (7)
S2 <sup>ii</sup> —K1—O1—C4	−43.2 (2)	O2—K1—O2 <sup>iv</sup> —K1 <sup>iv</sup>	−159.92 (8)
S2 <sup>ii</sup> —K1—O1—K1 <sup>iv</sup>	−154.62 (6)	K1 <sup>v</sup> —S1—C1—S2	39.50 (18)
O1 <sup>iii</sup> —K1—O1—C3	−10.0 (3)	K1 <sup>v</sup> —S1—C1—N1	−140.0 (2)
O1 <sup>iii</sup> —K1—O1—C4	−149.2 (2)	K1 <sup>ii</sup> —S1—C1—S2	−36.69 (18)
O1 <sup>iii</sup> —K1—O1—K1 <sup>iv</sup>	99.41 (13)	K1 <sup>ii</sup> —S1—C1—N1	143.8 (2)
O2 <sup>iv</sup> —K1—O1—C3	−157.5 (2)	K1 <sup>v</sup> —S2—C1—S1	−38.31 (18)
O2 <sup>iv</sup> —K1—O1—C4	63.4 (2)	K1 <sup>v</sup> —S2—C1—N1	141.2 (3)
O2 <sup>iv</sup> —K1—O1—K1 <sup>iv</sup>	−48.08 (8)	K1 <sup>ii</sup> —S2—C1—S1	36.78 (18)

O1—K1—O2—K1 <sup>iii</sup>	−113.36 (12)	K1 <sup>ii</sup> —S2—C1—N1	−143.7 (2)
S1 <sup>i</sup> —K1—O2—K1 <sup>iii</sup>	93.11 (11)	K1—O1—C3—C2	−81.8 (3)
S2 <sup>i</sup> —K1—O2—K1 <sup>iii</sup>	129.81 (6)	C4—O1—C3—C2	60.9 (3)
S1 <sup>ii</sup> —K1—O2—K1 <sup>iii</sup>	−16.20 (6)	K1 <sup>iv</sup> —O1—C3—C2	−178.43 (19)
S2 <sup>ii</sup> —K1—O2—K1 <sup>iii</sup>	−60.59 (5)	K1—O1—C4—C5	82.8 (3)
O1 <sup>iii</sup> —K1—O2—K1 <sup>iii</sup>	51.11 (7)	C3—O1—C4—C5	−60.7 (3)
O2 <sup>iv</sup> —K1—O2—K1 <sup>iii</sup>	−165.83 (8)	K1 <sup>iv</sup> —O1—C4—C5	179.8 (2)
O1—K1—S1 <sup>i</sup> —C1 <sup>i</sup>	−136.36 (13)	C2—N1—C1—S1	−176.4 (2)
O1—K1—S1 <sup>i</sup> —K1 <sup>iv</sup>	−48.73 (6)	C2—N1—C1—S2	4.1 (4)
O2—K1—S1 <sup>i</sup> —C1 <sup>i</sup>	27.08 (16)	C5—N1—C1—S1	−8.1 (4)
O2—K1—S1 <sup>i</sup> —K1 <sup>iv</sup>	114.71 (11)	C5—N1—C1—S2	172.4 (2)
O1—K1—S2 <sup>i</sup> —C1 <sup>i</sup>	81.34 (12)	C1—N1—C2—C3	−140.7 (3)
O1—K1—S2 <sup>i</sup> —K1 <sup>iv</sup>	−7.73 (6)	C5—N1—C2—C3	49.8 (3)
O2—K1—S2 <sup>i</sup> —C1 <sup>i</sup>	−132.18 (12)	C1—N1—C5—C4	140.7 (3)
O2—K1—S2 <sup>i</sup> —K1 <sup>iv</sup>	138.75 (7)	C2—N1—C5—C4	−50.0 (3)
O1—K1—S1 <sup>ii</sup> —C1 <sup>ii</sup>	69.70 (12)	N1—C2—C3—O1	−55.7 (3)
O1—K1—S1 <sup>ii</sup> —K1 <sup>iii</sup>	157.81 (5)	O1—C4—C5—N1	55.5 (3)

Symmetry codes: (i)  $-x, y+1/2, -z+3/2$ ; (ii)  $-x, -y, -z+1$ ; (iii)  $x, -y+1/2, z-1/2$ ; (iv)  $x, -y+1/2, z+1/2$ ; (v)  $-x, y-1/2, -z+3/2$ .

#### Hydrogen-bond geometry ( $\text{\AA}$ , $^\circ$ )

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
O2—H1O $\cdots$ S1 <sup>vi</sup>	0.86 (4)	2.45 (4)	3.219 (3)	149 (3)
O2—H2O $\cdots$ S1 <sup>vii</sup>	0.85 (3)	2.87 (5)	3.462 (3)	129 (4)

Symmetry codes: (vi)  $-x-1, y+1/2, -z+1/2$ ; (vii)  $-x-1, -y, -z+1$ .